



(43) International Publication Date 12 July 2001 (12.07.2001)

PCT

(10) International Publication Number WO 01/49728 A2

(51) International Patent Classification7: C07K 14/435

229-0014 (JP). KIMURA, Tomoko [JP/JP]; 715, 2-9-1, Kohoku, Tsuchiura-shi, Ibaraki 300-0032 (JP).

(21) International Application Number: PCT/JP00/09359

(74) Agents: AOYAMA, Tamotsu et al.; AOYAMA & PART-NERS, IMP Building, 3-7, Shiromi I-chome, Chuo-ku, Osaka-shi, Osaka 540-0001 (JP).

(22) International Filing Date:

28 December 2000 (28.12.2000)

English (8

(25) Filing Language: En

(81) Designated States (national): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CR, CU, CZ, DE, DK, DM, DZ, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW.

(26) Publication Language:

English

(30) Priority Data:

 2000-585
 6 January 2000 (06.01.2000)
 JP

 2000-588
 6 January 2000 (06.01.2000)
 JP

 2000-2299
 11 January 2000 (11.01.2000)
 JP

 2000-26862
 3 February 2000 (03.02.2000)
 JP

 2000-58367
 3 March 2000 (03.03.2000)
 JP

(84) Designated States (regional): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, TT, LU, MC, NL, PT, SE, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).

(71) Applicants (for all designated States except US): PROTEGENE INC. [JP/JP]; 2-20-3, Naka-cho, Meguro-ku, Tokyo 153-0065 (JP). SAGAMI CHEMICAL RESEARCH CENTER [JP/JP]; 4-1, Nishi-Ohnuma 4-chome, Sagamihara-shi, Kanagawa 229-0012 (JP).

Published:

 Without international search report and to be republished upon receipt of that report.

(72) Inventors; and

(75) Inventors/Applicants (for US only): KATO, Seishi [JP/JP]; 3-46-50, Wakamatsu, Sagamihara-shi, Kanagawa

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

1/49728 A2

(54) Title: HUMAN PROTEINS HAVING HYDROPHOBIC DOMAINS AND DNAs ENCODING THESE PROTEINS

(57) Abstract: The present invention provides human proteins having hydrophobic domains, DNAs encoding these proteins, expression vectors for these DNAs, transformed eukaryotic cells expressing these DNAs and antibodies directed to these proteins.

WO 01/49728 PCT/JP00/09359

DESCRIPTION

Human Proteins Having Hydrophobic Domains and DNAs Encoding These Proteins

5

10

15

20

TECHNICAL FIELD

The present invention relates to human proteins having hydrophobic domains, DNAs encoding these proteins, cells eukaryotic DNAs, expression vectors for these expressing these DNAs and antibodies directed to these proteins. The proteins of the present invention can be employed as pharmaceuticals or as antigens for preparing antibodies directed to these proteins. The human cDNAs of the present invention can be utilized as probes for genetic diagnosis and gene sources for gene therapy. Furthermore, the cDNAs can be utilized as gene sources for producing the proteins encoded by these cDNAs in large quantities. Cells into which these genes are introduced to express secretory proteins or membrane proteins in large quantities can be utilized for detection of the corresponding receptors or ligands, screening of novel small molecule pharmaceuticals and the like. The antibodies of the present invention can be utilized for the detection, quantification, purification and the like of the proteins of the present invention.

10

15

20

25

BACKGROUND ART

Cells secrete many proteins extracellularly. These secretory proteins play important roles in the proliferation induction, the material differentiation the control, transport, the biophylaxis, and the like of the cells. Unlike intracellular proteins, the secretory proteins exert their actions outside the cells. Therefore, they can be administered in the intracorporeal manner such as the they possess drip, and the or injection potentialities as pharmaceuticals. In fact, a number of human secretory proteins such as interferons, interleukins, erythropoietin, thrombolytic agents and the like pharmaceuticals. In addition, employed as currently secretory proteins other than those described above are undergoing clinical trials for developing their use as pharmaceuticals. It is believed that the human cells produce many unknown secretory proteins. Availability of these secretory proteins as well as genes encoding them is expected to lead to development of novel pharmaceuticals utilizing them.

On the other hand, membrane proteins play important roles, as signal receptors, ion channels, transporters and the like, in the material transport and the signal transduction through the cell membrane. Examples thereof include receptors for various cytokines, ion

channels for the sodium ion, the potassium ion, the chloride ion and the like, transporters for saccharides, amino acids and the like. The genes for many of them have already been cloned. It has been clarified that abnormalities in these membrane proteins are involved in a number of previously cryptogenic diseases. Therefore, discovery of a new membrane protein is expected to lead to elucidation of the causes of many diseases, and isolation of new genes encoding the membrane proteins has been desired.

.5

10

15

20

25

Heretofore, due to difficulty in the purification from human cells, many of these secretory proteins and membrane proteins have been isolated by genetic approaches. A general method is the so-called expression cloning method, in which a cDNA library is introduced into eukaryotic cells to express cDNAs, and the cells secreting, or expressing on the surface of membrane, the protein having the activity of interest are then screened. However, only genes for proteins with known functions can be cloned by using this method.

In general, a secretory protein or a membrane protein possesses at least one hydrophobic domain within the protein. After synthesis on ribosomes, such domain works as a secretory signal or remains in the phospholipid membrane to be entrapped in the membrane. Accordingly, if the existence of a highly hydrophobic domain is observed in the amino acid sequence of a protein encoded by a cDNA when the

whole base sequence of the full-length cDNA is determined, it is considered that the cDNA encodes a secretory protein or a membrane protein.

5 OBJECTS OF INVENTION

10

The main object of the present invention is to provide novel human proteins having hydrophobic domains, DNAs coding for these proteins, expression vectors for these DNAs, transformed eucaryotic cells that are capable of expressing these DNAs and antibodies directed to these proteins.

SUMMARY OF INVENTION

As the result of intensive studies, the present inventors have successfully cloned cDNAs encoding proteins 15 having hydrophobic domains from the human full-length cDNA bank, thereby completing the present invention. Thus, the human protein invention provides а hydrophobic domain(s), namely a protein comprising any one of amino acid sequences selected from the group consisting 20 of SEQ ID NOS: 1 to 10, 31 to 40, 61 to 70, 91 to 100 and 121 to 130. Moreover, the present invention provides a DNA encoding said protein, exemplified by a cDNA comprising any one of base sequences selected from the group consisting of SEQ ID NOS: 11 to 30, 41 to 60, 71 to 90, 101 to 120 and 131 25

to 150, an expression vector that is capable of expressing said DNA by in vitro translation or in eukaryotic cells, a transformed eukaryotic cell that is capable of expressing said DNA and of producing said protein, and an antibody directed to said protein.

This object as well as other objects and advantages of the present invention will become apparent to those skilled in the art from the following description with reference to the accompanying drawings.

10

15

20

5

BRIEF DESCRIPTION OF DRAWINGS

Figure 1: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP03613.

Figure 2: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP03700.

Figure 3: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP03935.

Figure 4: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP10755.

Figure 5: A figure depicting the
25 hydrophobicity/hydrophilicity profile of the protein

10

15

20

25

encoded by clone HP10760.

Figure 6: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP10764.

Figure 7: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP10768.

Figure 8: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP10769.

Figure 9: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP10784.

Figure 10:A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP10786.

Figure 11:A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP03727.

Figure 12:A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP03801.

Figure 13:A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP03883.

the depicting figure Figure 14: A hydrophobicity/hydrophilicity profile of the protein encoded by clone HP03913. depicting the figure Figure 15: A hydrophobicity/hydrophilicity profile of the protein 5 encoded by clone HP10753. the depicting figure Figure 16: A hydrophobicity/hydrophilicity profile protein of the encoded by clone HP10758. the depicting Figure 17: A figure 10 hydrophobicity/hydrophilicity profile protein of the encoded by clone HP10771. depicting the figure Figure 18: A hydrophobicity/hydrophilicity profile protein the of encoded by clone HP10778. 15 the depicting figure Figure 19: A hydrophobicity/hydrophilicity profile of protein the encoded by clone HP10781. the depicting figure Figure 20:A hydrophobicity/hydrophilicity profile of protein the 20 encoded by clone HP10785. figure depicting the Figure 21:A protein hydrophobicity/hydrophilicity profile of the encoded by clone HP03878. the figure depicting Figure 22:A 25

15

20

hydrophobicity/hydrophilicity profile of the protein encoded by clone HP03884.

Figure 23:A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP03934.

Figure 24: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP03949.

Figure 25: A figure depicting the

10 hydrophobicity/hydrophilicity profile of the protein

encoded by clone HP03959.

Figure 26: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP03983.

Figure 27: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP10745.

Figure 28: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP10775.

Figure 29: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP10782.

Figure 30:A figure depicting the hydrophobicity/hydrophilicity profile of the protein.

the figure depicting Figure 31:A hydrophobicity/hydrophilicity profile of the protein encoded by clone HP03977. figure depicting the Figure 32:A hydrophobicity/hydrophilicity profile of the protein 5 encoded by clone HP10649. depicting the Figure 33:A figure hydrophobicity/hydrophilicity profile of the protein encoded by clone HP10779. depicting the figure Figure 34: A 10 hydrophobicity/hydrophilicity profile of protein the encoded by clone HP10790. the depicting figure Figure 35: A hydrophobicity/hydrophilicity profile of the protein encoded by clone HP10793. 15 figure depicting the Figure 36: A hydrophobicity/hydrophilicity profile of protein the encoded by clone HP10794. Figure 37: A figure depicting the hydrophobicity/hydrophilicity profile protein of. the 20 encoded by clone HP10797. the figure depicting Figure 38: A hydrophobicity/hydrophilicity profile of protein the encoded by clone HP10798. figure depicting the Figure 39: A 25

15

20

hydrophobicity/hydrophilicity profile of the protein encoded by clone HP10800.

Figure 40:A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP10801.

Figure 41:A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP03596.

Figure 42:A figure depicting the 10 hydrophobicity/hydrophilicity profile of the protein encoded by clone HP03882.

Figure 43:A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP03903.

Figure 44: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP03974.

Figure 45: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP03978.

Figure 46: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP10735.

Figure 47: A figure depicting the
25 hydrophobicity/hydrophilicity profile of the protein

10

15

20

25

encoded by clone HP10750.

Figure 48: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP10777.

Figure 49: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP10780.

Figure 50:A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP10795.

DETAILED DESCRIPTION OF THE INVENTION

obtained, for example, by a method for isolating proteins from human organs, cell lines or the like, a method for preparing peptides by the chemical synthesis based on the amino acid sequence of the present invention, or a method for producing proteins by the recombinant DNA technology using the DNAs encoding the hydrophobic domains of the present invention. Among these, the method for producing proteins by the recombinant DNA technology is preferably employed. For example, the proteins can be expressed in vitro by preparing an RNA by in vitro transcription from a vector having the cDNA of the present invention, and then carrying out in vitro translation using this RNA as a

region into a suitable expression vector by the method known in the art may lead to expression of the encoded protein in large quantities in prokaryotic cells such as *Escherichia* coli and *Bacillus* subtilis, or eukaryotic cells such as yeasts, insect cells and mammalian cells.

5

10

15

20

25

In the case where the protein of the present invention is produced by expressing the DNA by in vitro translation, the protein of the present invention can be produced in vitro by incorporating the translated region of this cDNA into a vector having an RNA polymerase promoter, and then adding the vector to an in vitro translation system such as a rabbit reticulocyte lysate or a wheat germ extract, which contains an RNA polymerase corresponding to the promoter. The RNA polymerase promoters are exemplified by T7, T3, SP6 and the like. The vectors containing promoters for these RNA polymerases are exemplified by pKA1, pCDM8, pT3/T7 18, pT7/3 19, pBluescript II and the like. Furthermore, the protein of the present invention can be expressed in the secreted form or the form incorporated in the microsome membrane when a canine pancreas microsome or the like is added to the reaction system.

In the case where the protein of the present invention is produced by expressing the DNA in a microorganism such as *Escherichia coli*, a recombinant

10

15

20

25

expression vector in which the translated region of the cDNA of the present invention is incorporated into an expression vector having an origin which is capable of replicating in the microorganism, a promoter, a ribosome-binding site, a cDNA-cloning site, a terminator and the like is constructed. After transformation of the host cells with this expression vector, the resulting transformant is cultured. Thus, the protein encoded by the cDNA can be produced in large quantities in the microorganism. In this case, a protein fragment containing any translated region can be obtained by adding an initiation codon and a termination codon in front of and behind the selected translated region and expressing the protein. Alternatively, the protein can be expressed as a fusion protein with another protein. Only the portion of the protein encoded by the cDNA can be obtained by cleaving this fusion protein with a suitable protease. The expression vectors for Escherichia coli are exemplified by the pUC series, pBluescript II, the pET expression system, the pGEX expression system and the like.

In the case where the protein of the present invention is produced by expressing the DNA in eukaryotic cells, the protein of the present invention can be produced as a secretory protein, or as a membrane protein on the surface of cell membrane, by incorporating the translated region of the cDNA into an expression vector for eukaryotic

10

15

20

25

cells that has a promoter, a splicing region, a poly(A) addition site and the like, and then introducing the vector into the eukaryotic cells. The expression vectors are exemplified by pKA1, pED6dpc2, pCDM8, pSVK3, pMSG, pSVL, pBK-CMV, pBK-RSV, EBV vectors, pRS, pYES2 and the like. Examples of eukaryotic cells to be used in general include mammalian cultured cells such as monkey kidney COS7 cells and Chinese hamster ovary CHO cells, budding yeasts, fission yeasts, silkworm cells, and Xenopus oocytes. Any eukaryotic cells may be used as long as they are capable of expressing the proteins of the present invention. The expression vector can be introduced into the eukaryotic cells by using a method known in the art such as the electroporation method, the calcium phosphate method, the liposome method and the DEAE-dextran method.

After the protein of the present invention is expressed in prokaryotic cells or eukaryotic cells, the protein of interest can be isolated and purified from the culture by a combination of separation procedures known in the art. Examples of the separation procedures include treatment with a denaturing agent such as urea or a detergent, sonication, enzymatic digestion, salting-out or dialysis, centrifugation, precipitation, solvent filtration, SDS-PAGE, isoelectric ultrafiltration, gel chromatography, hydrophobic focusing, ion-exchange

10

15

20

25

PCT/JP00/09359

chromatography, affinity chromatography and reverse phase chromatography.

The proteins of the present invention also include peptide fragments (of 5 amino acid residues or more) containing any partial amino acid sequences in the amino acid sequences represented by SEQ ID NOS: 1 to 10, 31 to 40, 61 to 70, 91 to 100 and 121 to 130. These peptide fragments can be utilized as antigens for preparation of antibodies. Among the proteins of the present invention, those having the signal sequences are secreted in the form of mature proteins after the signal sequences are removed. Therefore, these mature proteins shall come within the scope of the protein of the present invention. The N-terminal amino acid sequences of the mature proteins can be easily determined by using the method for the determination of cleavage site of a signal sequence [JP-A 8-187100]. Furthermore, some membrane proteins undergo the processing on the cell surface to be converted to the secreted forms. Such proteins or peptides in the secreted forms shall also come within the scope of the protein of the present invention. In the case where sugar chain-binding sites are present in the amino acid sequences of the proteins, expression of the proteins in appropriate eukaryotic cells affords the proteins to which sugar chains are added. Accordingly, such proteins or peptides to which sugar chains are added shall also come WO 01/49728 PCT/JP00/09359

within the scope of the protein of the present invention.

. 5

10

15

20

25

The DNAs of the present invention include all the DNAs encoding the above-mentioned proteins. These DNAs can be obtained by using a method for chemical synthesis, a method for cDNA cloning and the like.

The cDNAs of the present invention can be cloned, for example, from cDNA libraries derived from the human cells. The cDNAs are synthesized by using poly(A) + RNAs extracted from human cells as templates. The human cells may be cells delivered from the human body, for example, by the operation or may be the cultured cells. The cDNAs can be synthesized by using any method such as the Okayama-Berg method [Okayama, H. and Berg, P., Mol. Cell. Biol. 2: 161-170 (1982)], the Gubler-Hoffman method [Gubler, U. and Hoffman, J., Gene 25: 263-269 (1983)] and the like. However, it is desirable to use the capping method [Kato, S. et al., Gene 150: 243-250 (1994)], as exemplified in Examples, in order to obtain a full-length clone in an effective manner. In addition, commercially available human cDNA libraries can be utilized. The cDNAs of the present invention can be by synthesizing **cDNA** libraries the from cloned oligonucleotide on the basis of base sequences of any portion in the cDNA of the present invention and screening the cDNA libraries using this oligonucleotide as a probe for colony or plaque hybridization according to a method known

10

15

in the art. In addition, the cDNA fragments of the present invention can be prepared from an mRNA isolated from human cells by the RT-PCR method in which oligonucleotides which hybridize with both termini of the cDNA fragment of interest are synthesized, which are then used as the primers.

The cDNAs of the present invention are characterized in that they comprise any one of the base sequences represented by SEQ ID NOS: 11 to 20, 41 to 50, 71 to 80, 101 to 110 and 131 to 140 or the base sequences represented by SEQ ID NOS: 21 to 30, 51 to 60, 81 to 90, 111 to 120 and 141 to 150. Table 1 summarizes the clone number (HP number), the cells from which the cDNA clone was obtained, the total number of bases of the cDNA, and the number of the amino acid residues of the encoded protein, for each of the cDNAs.

Table 1

<u></u>													
		Sequence No.		HP No.	Cell	Number	Number of						
						of	amino						
						bases	acids						
	1,	11,	21	HP03613	Kidney	2865	578						
	2,	12,	22	нр03700	Kidney	3323	243						
	3,	13,	23	HP03935	Kidney	1585	461						
	4,	14,	24	HP10755	Kidney	2122	647						
	5,	15,	25	HP10760	Kidney	1775	446						
	6,	16,	26	HP10764	Kidney	1372	197						
	7,	17,	27	HP10768	Kidney	2074	540						
	8,	18,	28	HP10769	Kidney	2252	442						
	9,	19,	29	HP10784	Kidney	1461	262						
	10,	20,	30	HP10786	Kidney	1122	152						
•	31,	41,	51	нр03727	Kidney	1617	335						
	32,	42,	52	HP03801	Umbilical cord blood	1749	208						
	33,	43,	53	HP03883	Kidney	1402	406						
	34,	44,	54	HP03913	Kidney	2474	618						
	35,	45,	55	HP10753	Umbilical cord blood	3296	208						
	36,	46,	56	HP10758	Kidney	1818	502						
	37,	47,	57	HP10771	Kidney	1646	336						
	38,	48,	58	HP10778	Kidney	1416	340						
	39,	49,	59	HP10781	Kidney	1927	223						
	40,	50,	60	HP10785	Kidney	1419	309						
	61,	71,	81	HP03878	Kidney	2016	599						
	62,	72,	82	HP03884	Kidney	1446	81						
	63,	73,	83	HP03934	Kidney	2467	654						
	64,	74,	84	HP03949	Kidney	1450	390						
_	65,	75,	85	HP03959	Kidney	1897	452						
			•										

WO 01/49728 PCT/JP00/09359

invention can be easily obtained by screening the cDNA libraries constructed from the human cell lines or human tissues utilized in the present invention using an oligonucleotide probe synthesized on the basis of the base sequence of the cDNA provided in any one of SEQ ID NOS: 11 to 30, 41 to 60, 71 to 90, 101 to 120 and 131 to 150.

5

10

15

20

25

In general, the polymorphism due to the individual differences is frequently observed in human genes. Accordingly, any cDNA in which one or plural nucleotides are added, deleted and/or substituted with other nucleotides in SEQ ID NOS: 11 to 30, 41 to 60, 71 to 90, 101 to 120 and 131 to 150 shall come within the scope of the present invention.

Similarly, any protein in which one or plural amino acids are added, deleted and/or substituted with other amino acids resulting from the above-mentioned changes shall come within the scope of the present invention, as long as the protein possesses the activity of the protein having any one of the amino acid sequences represented by SEQ ID NOS: 1 to 10, 31 to 40, 61 to 70, 91 to 100 and 121 to 130.

The cDNAs of the present invention also include cDNA fragments (of 10 bp or more) containing any partial base sequence in the base sequences represented by SEQ ID NOS: 11 to 20, 41 to 50, 71 to 80, 101 to 110 and 131 to 140 or in the base sequences represented by SEQ ID NOS: 21 to 30, 51 to 60, 81 to 90, 111 to 120 and 141 to 150. Also, DNA

WO 01/49728 PCT/JP00/09359

fragments each consisting of a sense strand and an antisense strand shall come within this scope. These DNA fragments can be utilized as the probes for the genetic diagnosis.

5

10

15

20

25

obtained from a serum after immunizing an animal using the protein of the present invention as an antigen. A peptide that is chemically synthesized based on the amino acid sequence of the present invention and a protein expressed in eukaryotic or prokaryotic cells can be used as an antigen. Alternatively, an antibody can be prepared by introducing the above-mentioned expression vector for eukaryotic cells into the muscle or the skin of an animal by injection or by using a gene gun and then collecting a serum therefrom [JP-A 7-313187]. Animals that can be used include a mouse, a rat, a rabbit, a goat, a chicken and the like. A monoclonal antibody directed to the protein of the present invention can be produced by fusing B cells collected from the spleen of the immunized animal with myelomas to generate hybridomas.

In addition to the activities and uses described above, the polynucleotides and proteins of the present invention may exhibit one or more of the uses or biological activities (including those associated with assays cited herein) identified below. Uses or activities described for proteins of the present invention may be provided by

10

15

20

25

administration or use of such proteins or by administration or use of polynucleotides encoding such proteins (such as, for example, in gene therapies or vectors suitable for introduction of DNA).

Research Uses and Utilities

The polynucleotides provided by the present invention can be used by the research community for various purposes. The polynucleotides can be used to express for analysis, characterization recombinant protein therapeutic use; as markers for tissues in which the corresponding protein is preferentially expressed (either constitutively or at a particular stage differentiation or development or in disease states); as molecular weight markers on Southern gels; as chromosome markers or tags (when labeled) to identify chromosomes or to map related gene positions; to compare with endogenous DNA patients to identify potential sequences in disorders; as probes to hybridize and thus discover novel, related DNA sequences; as a source of information to derive PCR primers for genetic fingerprinting; as a probe to "subtract-out" known sequences in the process of discovering other novel polynucleotides; for selecting and making oligomers for attachment to a "gene chip" or other support, including for examination of expression patterns; to raise anti-protein antibodies using DNA immunization techniques;

10

15

20

25

and as an antigen to raise anti-DNA antibodies or elicit another immune response. Where the polynucleotide encodes a protein which binds or potentially binds to another protein (such as, for example, in a receptor-ligand interaction), the polynucleotide can also be used in interaction trap assays (such as, for example, that described in Gyuris et al., Cell 75:791-803 (1993)) to identify polynucleotides encoding the other protein with which binding occurs or to identify inhibitors of the binding interaction.

The proteins provided by the present invention can similarly be used in assay to determine biological activity, including in a panel of multiple proteins for highthroughput screening; to raise antibodies or to elicit another immune response; as a reagent (including the labeled reagent) in assays designed to quantitatively determine levels of the protein (or its receptor) in biological fluids; as markers for tissues in which the corresponding protein is preferentially expressed (either constitutively or at a particular stage of tissue differentiation or development or in a disease state); and, of course, to isolate correlative receptors or ligands. Where the protein binds or potentially binds to another protein (such as, for example, in a receptor-ligand interaction), the protein can be used to identify the other protein with which binding occurs or to identify inhibitors of the binding interaction.

10

15

20

25

base sequences of the present cDNA has revealed the registration of sequences that shared a homology of 90% or more (for example, Accession No. AA42490) among ESTs. However, since they are partial sequences, it can not be judged whether or not they encode the same protein as the protein of the present invention.

<HP10760> (SEQ ID NOS: 5, 15, and 25)

Determination of the whole base sequence of the cDNA insert of clone HP10760 obtained from cDNA library of human kidney revealed the structure consisting of a 61-bp 5'-untranslated region, a 1341-bp ORF, and a 373-bp 3'untranslated region. The ORF encodes a protein consisting of 446 amino acid residues and there existed a putative secretory signal at the N-terminus. Figure 5 depicts the hydrophobicity/hydrophilicity profile, obtained by the Kytethe present protein. In vitro Doolittle method, of translation resulted in formation of a translation product of 48 kDa that was somewhat smaller than the molecular weight of 49,468 predicted from the ORF. In this case, the addition of a microsome led to the formation of a product of 50 kDa. In addition, there exists in the amino acid sequence of this protein two sites at which N-glycosylation may occur (Asn-Ala-Thr at position 144 and Asn-Ile-Ser at position 243). Application of the (-3,-1) rule, a method for predicting the cleavage site of the secretory signal sequence, allows to expect that the mature protein starts from glutamic acid at position 27.

The search of the protein database using the amino acid sequence of the present protein revealed that the protein was similar to human 25 kDa trypsin inhibitor (Accession No. BAA25066). Table 6 shows the comparison between amino acid sequences of the human protein of the present invention (HP) and human 25 kDa trypsin inhibitor (TI). Therein, the marks of -, *, and . represent a gap, an amino acid residue identical with that of the protein of the present invention, and an amino acid residue similar to that of the protein of the present invention, respectively. The both proteins shared a homology of 33.5% in the intermediate region of 185 amino acid residues.

15

10

5

Table 6

HР

MLHPETSPGRGHLLAVLLALLGTAWAEVWPPQLQEQAPMAG

20 TI MIAISAVSSALLFSLLCEASTVVLLNSTDSSPPTNNFTDIEAALKAQLDSADIPKARRKR

15

20

25

	HP	RTLQVGWNMQLLPAGLASFVEVVSLWFAEGQRYSHA-AGECAR-NATCTHYTQL
		* * * * * * ******
•	ΤI	RFLGQNLSVRTGRYRSILQLVKPWYDEVKDYAFPYPQDCNPRCPMRCFGPMCTHYTQM
		DAE WCAVCDCCANWEVAICUTTTDVVVCAWCCLC
5	HP	VWATSSQLGCGRHLCSAGQA—AI——EAF-VCAYSPGGNWEVNGKTIIPYKKGAWCSLC
		***** * * * * * * * * * * * * * * * * *
	TI	VWATSNRIGCAIHTCQNMNVWGSVWRRAVYLVCNYAPKGNWIGEAPYKVGVPCSSC
	HP	TASVSGCFKAWDHAGGLCEVPRNPCRMSCQNHGRLNISTCHCHCPPGYTGRYCQVRCSLQ

TI PPSYGGSCTDNLCFPGVTSNYLYWFK

..*.*

The search of the GenBank using the base sequences of the present cDNA has revealed the registration of sequences that shared a homology of 90% or more (for example, Accession No. AI792411) among ESTs. However, since they are partial sequences, it can not be judged whether or not they encode the same protein as the protein of the present invention.

<HP10764> (SEQ ID NOS: 6, 16, and 26)

Determination of the whole base sequence of the cDNA insert of clone HP10764 obtained from cDNA library of human kidney revealed the structure consisting of a 326-bp 5'-untranslated region, a 594-bp ORF, and a 452-bp 3'-untranslated region. The ORF encodes a protein consisting of

WO 01/49728 PCT/JP00/09359

12 /346

				500					505					510		
	Leu	Ala	Val	Gly	Ala	Ala	Phe	Ala	ser	Ser	Trp	Lys	Thr	Gly	Leu	Ala
			515					520					525			
	Thr	Ser	Leu	Ala	Val	Phe	Суз	His	Glu	Leu	Pro	His	Glu	Leu	Gly	Asp
5		530					535					540				
	Phe	Ala	Ala	Leu	Leu	His	Ala	Gly	Leu	Ser	Val	Arg	Gln	Ala	Leu	Leu
	545					550					555					560
	Leu	Asn	Leu	Ala	Ser	Ala	Leu	Thr	Ala	Phe	Ala	GLy	Leu	Tyr	Val	Ala
					565					570					575	
10	Leu	Ala	Val	Gly	Val	Ser	Glu	Glu	Ser	Glu	Ala	Trp	Ile	Leu	Ala	Val
				580					585					590		
	Ala	Thr	Gly	Leu	Phe	Leu	Tyr	Val	Ala	Leu	Cys	Asp	Met	Leu	Pro	Ala
			595					600					605			
	Met	Leu	Lys	Val	Arg	Asp	Pro	Arg	Pro	Trp	Leu	Leu	Phe	Leu	Leu	His
15		610	1				615					620				
	Asn	Val	Gly	Leu	Leu	Gly	Gly	Trp	Thr	Val	Leu	Leu	Leu	Leu	Ser	Leu
	625	•				630					635					640
	Tyr	Glu	Asp	Asp	Ile	Thr	Phe	:								
					645	•										
20																
	<21	10> 5	5													
	<21	1> 4	146													
	<21	L2> 1	PRT										,			
	<23	L3> 1	Homo	sap	Lens											

25 <400> 5

WO 01/49728 PCT/JP00/09359

13 /346

	Met	Leu	His	Pro	Glu	Thr	Ser	Pro	Gly	Arg	Gly	His	Leu	Leu	Ala	Val
	1				5					10					15	
	Leu	Leu	Ala	Leu	Leu	Gly	Thr	Ala	Trp	Ala	Glu	Val	Trp	Pro	Pro	Gln
				20					25					30		
5	Leu	Gln	Glu	Gln	Ala	Pro	Met	Ala	Gly	Ala	Leu	Asn	Arg	Lys	Glu	Ser
			35					40					45			
	Phe	Leu	Leu	Leu	Ser	Leu	His	Asn	Arg	Leu	Arg	Ser	Trp	Val	Gln	Pro
		50					55					60				
	Pro	Ala	Ala	Asp	Met	Arg	Arg	Leu	Asp	Trp	Ser	Asp	Ser	Leu	Ala	Gln
10	65					70					75					80
	Leu	Ala	Glņ	Ala	Arg	Ala	Ala	Leu	Cys	Gly	Ile	Pro	Thr	Pro	Ser	Leu
					85					90					95	
	Ala	Ser	Gly	Leu	Trp	Arg	Thr	Leu	Gln	Val	Gly	Trp	Asn	Met	Gln	Leu
				100					105					110		
15	Leu	Pro	Ala	Gly	Leu	Ala	Ser	Phe	Val	Glu	Val	Val	Ser	Leu	Trp	Phe
			115					120					125			
	Ala	Glu	Gly	Gln	Arg	Tyr	Ser	His	Ala	Ala	Gly	Glu	Cys	Ala	Arg	Asn
		130					135					140				
	Ala	Thr	Cys	Thr	His	Tyr	Thr	Gln	Leu	Val	Trp	Ala	Thr	Ser	Ser	Gln
20	145					150					155					160
	Leu	Gly	Cys	Gly	Arg	His	Leu	Cys	Ser	Ala	Gly	Gln	Ala	Ala	Ile	Glu
					165					170	!				175	
	Ala	Phe	Val	. Cys	Ala	Tyr	Ser	Pro	Gly	Gly	Asn	Trp	G1u	Val	Asn	Gly
				180	i				185	,				190	i	
25	Lys	Thr	: Ile	: Ile	Pro	Tyr	Lys	Lys	Gly	, Ala	Trp	Cys	Ser	Leu	Cys	Thr

14 /346

			195					200					205			
	Ala	Ser	Val	Ser	Gly	Суз	Phe	Lys	Ala	Trp	Asp	His	Ala	Gly	Gly	Leu
		210					215				•	220				
	Cys	Glu	Val	Pro	Arg	Asn	Pro	Cys	Arg	Met	Ser	Cys	Gln	Asn	His	Gly
5	225					230					235					240
	Arg	Leu	Asn	Ile	Ser	Thr	Cys	His	Cys	Hìs	Cys	Pro	Pro	Gly	Tyr	Thr
					245					250					255	
	Gly	Arg	Tyr	Суз	Gln	Val	Arg	Cys	Ser	Leu	Gln	Cys	Val	His	Gly	Arg
				260					265					270		
10	Phe	Arg	Glu	Glu	Glu	Суз	Ser	Cys	Val	Суѕ	Asp	Ile	Gly	Tyr	Gly	Gly
			275					280					285			
	Ala	Gln	Cys	Ala	Thr	Lys	Val	His	Phe	Pro	Phe	His	Thr	Cys	Asp	Leu
		290					295					300				
	Arg	Ile	Asp	Gly	Asp	Суѕ	Phe	Met	Val	Ser	Ser	Glu	Ala	Asp	Thr	Туг
15	305					310					315					320
	Tyr	Arg	Ala	Arg	Met	Lys	Cys	Gln	Arg	Lys	Gly	Gly	Val	Leu	Ala	Gln
					325	,				330					335	
	Ile	Lys	Ser	Gln	Lys	Val	Gln	Asp	Ile	Leu	Ala	Phe	Tyr	Leu	Gly	Arg
				340)				345	,				350	ı	
20	Leu	Glu	Thr	Thr	Asn	Glu	Val	Ile	Asp	Ser	Asp	Phe	Glu	Thr	Arg	Asr
			355	5				360	ı				365	i		
	Phe	Trp	ıle	e Gly	, Leu	ı Thr	Tyr	Lys	Thr	Ala	Lys	asp	Sei	: Phe	Arg	Tr
		370)				375	;				380)			
	Ala	a Thi	c Gly	/ Glu	ı His	s Glr	n Ala	. Phe	Thi	: Ser	Phe	ala	a Phe	e Gly	Glr	n Pro
25	381	;				390)				395	5				400

PCT/JP00/09359 WO 01/49728

15 /346

Asp Asn His Gly Phe Gly Asn Cys Val Glu Leu Gln Ala Ser Ala Ala Phe Asn Trp Asn Asn Gln Arg Cys Lys Thr Arg Asn Arg Tyr Ile Cys Gln Phe Ala Gln Glu His Ile Ser Arg Trp Gly Pro Gly Ser <210> 6 <211> 197 <212> PRT <213> Homo sapiens <400> 6 Met Pro Pro Ala Gly Leu Arg Arg Ala Ala Pro Leu Thr Ala Ile Ala Leu Leu Val Leu Gly Ala Pro Leu Val Leu Ala Gly Glu Asp Cys Leu Trp Tyr Leu Asp Arg Asn Gly Ser Trp His Pro Gly Phe Asn Cys Glu Phe Phe Thr Phe Cys Cys Gly Thr Cys Tyr His Arg Tyr Cys Cys Arg Asp Leu Thr Leu Leu Ile Thr Glu Arg Gln Gln Lys His Cys Leu Ala Phe Ser Pro Lys Thr Ile Ala Gly Ile Ala Ser Ala Val Ile Leu Phe Val Ala Val Val Ala Thr Thr Ile Cys Cys Phe Leu Cys Ser Cys Cys

tggcgcctcc tggctatgct ggccgggctc tacgccttct tcctgtttga gaacctcttc 1260 aatctcctgc tgcccaggga cccggaggac ctggaggacg ggccctgcgg ccacagcagc 1320 catagecaeg ggggecaeag ceaeggtgtg teeetgeage tggcaeceag egageteegg 1380 cagcccaagc cccccacga gggctcccgc gcagacctgg tggcggagga gagcccggag 1440 ctgctgaacc ctgagcccag gagactgagc ccagagttga ggctactgcc ctatatgatc 1500 5 actotgggcg acgccgtgca caacttcgcc gacgggctgg ccgtgggcgc cgccttcgcg 1560 tecteetgga agaceggget ggeeaceteg etggeegtgt tetgeeacga gttgeeacae 1620 gagetggggg acttegeege ettgetgeac geggggetgt eegtgegeea ageactgetg 1680 ctgaacctgg cctccgcgct cacggccttc gctggtctct acgtggcact cgcggttgga 1740 gtcagcgagg agagcgaggc ctggatcctg gcagtggcca ccggcctgtt cctctacgta 1800 10 gcactetgeg acatgetece ggcgatgttg aaagtacggg accegeggee etggeteete 1860 ttcctgctgc acaacgtggg cctgctgggc ggctggaccg tcctgctgct gctgtccctg 1920 1944 tacgaggatg acatcacctt ctga

15 <210> 15

<211> 1341

<212> DNA

<213> Homo sapiens

<400> 15

atgctgcatc cagagacete ecetggeegg gggcatetee tggetgtget ectggceete 60

ettggcaceg cetgggeaga ggtgtggeea ececagetge aggageagge tecgatggee 120

ggagecetga acaggaagga gagtttettg etecteteee tgeacaaceg ectgegeage 180

tgggteeage ecectgegge tgacatgegg aggetggact ggagtgacag ectggeecaa 240

etggeteaag ecagggeage ectetgtgga ateceaacee egageetgge gteeggeetg 300

tggegeacee tgeaagtggg etggaacatg eagetgetge eegeggett ggegteettt 360

gttgaagtgg tcagcctatg gtttgcagag gggcagcggt acagccacgc ggcaggagag 420 tgtgctcgca acgccacctg cacccactac acgcagctcg tgtgggccac ctcaagccag 480 ctgggctgtg ggcggcacct gtgctctgca ggccaggcag cgatagaagc ctttgtctgt 540 gcctactccc ccggaggcaa ctgggaggtc aacgggaaga caatcatccc ctataagaag 600 ggtgcctggt gttcgctctg cacagccagt gtctcaggct gcttcaaagc ctgggaccat 660 gcagggggc tctgtgaggt ccccaggaat ccttgtcgca tgagctgcca gaaccatgga 720 cgtctcaaca tcagcacctg ccactgccac tgtccccctg gctacacggg cagatactgc 780 caagtgaggt gcagcctgca gtgttgtgcac ggccggttcc gggaggagga gtgctcgtgc 840 gtctgtgaca tcggctacgg gggagcccag tgtgccacca aggtgcattt tcccttccac 900 acctgtgacc tgaggatcga cggagactgc ttcatggtgt cttcagaggc agacacctat 960 10 tacagagcca ggatgaaatg tcagaggaaa ggcggggtgc tggcccagat caagagccag 1020 aaagtgcagg acatcctcgc cttctatctg ggccgcctgg agaccaccaa cgaggtgatt 1080 gacagtgact tcgagaccag gaacttctgg atcgggctca cctacaagac cgccaaggac 1140 teetteeget gggccacagg ggagcaccag geetteacca gttttgeett tgggcageet 1200 gacaaccacg ggtttggcaa ctgcgtggag ctgcaggctt cagctgcctt caactggaac 1260 15 aaccageget gcaaaacceg aaaccgttac atctgccagt ttgcccagga gcacatctcc 1320 1341 cggtggggcc cagggtcctg a

<210> 16

20 <211> 594

25

<212> DNA

<213> Homo sapiens

<400> 16

atgccacccg cggggctccg ccgggccgcg ccgctcaccg caatcgctct gttggtgctg 60 ggggctcccc tggtgctggc cggcgaggac tgcctgtggt acctggaccg gaatggctcc 120 WO 01/49728 PCT/JP00/09359

54 /346

635 640 630 gag gat gac atc acc ttc tgataccctg ccctagtccc ccacctttga 2026 Glu Asp Asp Ile Thr Phe 645 cttaagatcc cacacctcac aaacctacag cccagaaacc agaagcccct atagaggccc 2086 5 2122 cagteceaac tecagtaaag acactettgt cettgg <210> 25 <211> 1775 10 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (62)..(1402) 15 <400> 25 aaaacaagcc gggtggctga gccaggctgt gcacggagtg cctgacgggc ccaacagacc 60 c atg ctg cat cca gag acc tcc cct ggc cgg ggg cat ctc ctg gct gtg 109 Met Leu His Pro Glu Thr Ser Pro Gly Arg Gly His Leu Leu Ala Val 10 15 5 1 ctc ctg gcc ctc ctt ggc acc gcc tgg gca gag gtg tgg cca ccc cag 157 20 Leu Leu Ala Leu Leu Gly Thr Ala Trp Ala Glu Val Trp Pro Pro Gln 30 25 20 ctg cag gag cag gct ccg atg gcc gga gcc ctg aac agg aag gag agt 205

25 35 40 45

Leu Gln Glu Gln Ala Pro Met Ala Gly Ala Leu Asn Arg Lys Glu Ser

55 /346

	ttc tt	g ct	cc o	ctc	tcc	ctg	cac	aac	cgc	ctg	cgc	agc	tgg	gtc	cag	ccc	253
	Phe Le	u Le	eu I	Leu	Ser	Leu	His	Asn	Arg	Leu	Arg	Ser	Trp	Val	Gln	Pro	
	5	0					55					60					
	cct gc	g g	ct (gac	atg	cgg	agg	ctg	gac	tgg	agt	gac	agc	ctg	gcc	caa	301
5	Pro Al	a A	la :	Asp	Met	Arg	Arg	Leu	Asp	Trp	Ser	Asp	Ser	Leu	Ala	Gln	
	65					70					75					80	
	ctg go	ct c	aa	gcc	agg	gca	gcc	ctc	tgt	gga	atc	cca	acc	ccg	agc	ctg	349
	Leu A	La G	ln	Ala	Arg	Ala	Ala	Leu	Cys	Gly	Ile	Pro	Thr	Pro	Ser	Leu	
					85					90					95		
10	gcg t	cc g	gc	ctg	tgg	cgc	acc	ctg	caa	gtg	ggc	tgg	aac	atg	cag	ctg	397
	Ala S	er G	Sly	Leu	Trp	Arg	Thr	Leu	Gln	Val	Gly	Trp	Asn	Met	: Gln	Leu	
				100					105	1				110)		
	ctg c	cc q	gcg	ggc	ttg	gcg	tcc	ttt	. gtt	gaa	gtg	gto	ago	cta	a tgg	, ttt	445
	Leu P	ro I	Ala	Gly	Leu	Ala	Ser	Phe	val	. Glu	ı Val	. Val	L Sei	. Lei	ı Try	Phe	٠
15		;	115					120)			•	125	5			
	gca g	jag (ggg	cag	g egg	, tac	ago	cac	gc	g gca	a gga	a gag	g tgi	t gc	t cgo	c aac	493
	Ala	Slu	Gly	Glr	n Arg	ј Туг	: Sei	c His	s Ala	a Ala	a Gly	y Gl	u Cy	s Al	a Ar	g Asn	
	1	130					135	5				14	0				
																c cag	541
20	Ala	Thr	Cys	s Thi	r Hi	з Ту	r Th	r Gl	n Le	u ĮVa	l Tr	p Al	a Th	r Se	r Se	r Gln	
	145					15					15					160	
																a gaa	
	Leu	Gly	Cys	s Gl	y Ar	g Hi	s Le	u Cy	s Se	r Al	a Gl	y G1	n Al	a Al	la Il	e Glu	
					16					17					17		
25	gcc	ttt	gt	c tg	ıt go	c ta	c to	c co	c gg	ga gg	gc aa	ac to	gg ga	ag gi	tc aa	ac ggg	637

56/346

	Ala	Phe	Val	Суз	Ala	Tyr	Ser	Pro	Gly	Gly	Asn	Trp	Glu	Val	Asn	Gly	
				180					185					190			
	aag	aca	atc	atc	ccc	tat	aag	aag	ggt	gcc	tgg	tgt	tcg	ctc	tgc	aca	, 685
	Lys	Thr	Ile	Ile	Pro	Tyr	Lys	Lys	Gly	Ala	Trp	Cys	Ser	Leu	Cys	Thr	
5			195					200					205				
	gcc	agt	gtc	tca	ggc	tgc	ttc	aaa	gcc	tgg	gac	cat	gca	ggg	ggg	ctc	733
	Ala	Ser	Val	Ser	Gly	Cys	Phe	Lys	Ala	Trp	Asp	His	Ala	Gly	Gly	Leu	
		210	1				215					220					
	tgt	gag	gto	ccc	agg	aat	cct	tgt	cgc	atg	agc	tgc	cag	aac	cat	gga :	781
10	Cys	Glu	ı Val	Pro	Arg	Asn	Pro	Cys	Arg	Met	Ser	Суз	Gln	Asn	His	Gly	
	225)				230	ı				235					240	
	cgt	cto	c aac	c ato	ago	acc	: tgc	cac	tgc	cac	tgt	ccc	cct	ggc	tac	cacg	829
	Arg	, Lei	ı Ası	n Ile	e Sei	Thr	Cys	His	s Cys	His	Суз	Pro	Pro	Gly	у Ту	r Thr	
					245	5				250)				25	5	
15	ggo	c ag	a ta	c tgo	c caa	a gto	g agg	, tg	c ago	c ctg	cag	g tgt	gt	g ca	c gg	c cgg	877
	Gl	y Ar	g Ty	r Cy	s Gl	n Vai	l Arg	д Су:	s Sei	r Leu	ı Glı	ı Cys	s Vai	l Hi:	s Gl	y Arg	
				26	0				26	5				27	0		
	tt	c cg	g ga	g ga	g ga	g tg	c to	g tg	c gt	c tgl	ga ga	c at	c gg	c ta	c gg	g gga	925
	Ph	e Ar	g Gl	u Gl	u Gl	u Cy	s Se	r Cy	s Va	1 Cy:	s As	p Il	e Gl	у Ту	r Gl	y Gly	•
20			27	'5				28	0				28	5			
	gc	c ca	ng to	gt go	c ac	c aa	g gt	g ca	t tt	t cc	c tt	с са	c ac	c to	ıt ga	ac cto	973
	Al	a G]	in Cy	/s Al	a Th	ır Ly	s Va	l Hi	s Ph	e Pr	o Ph	e Hi	s Th	ır Cy	rs As	sp Leu	1
		29	90				29	5				30	0				
	aç	gg at	tc ga	ac gg	ga ga	ac to	jc tt	.c at	g gt	g to	t to	a ga	ag go	ca ga	ac a	cc ta	1021
25	Άι	g I	le A	sp G	Ly As	sp Cy	ys Ph	e Me	et Va	al Se	r Se	er Gl	lu Al	la As	sp T	hr Ty:	r

tac aga gcc agg atg aaa tgt cag agg aaa ggc ggg gtg ctg gcc cag Tyr Arg Ala Arg Met Lys Cys Gln Arg Lys Gly Gly Val Leu Ala Gln atc aag agc cag aaa gtg cag gac atc ctc gcc ttc tat ctg ggc cgc Ile Lys Ser Gln Lys Val Gln Asp Ile Leu Ala Phe Tyr Leu Gly Arg ctg gag acc acc aac gag gtg att gac agt gac ttc gag acc agg aac Leu Glu Thr Thr Asn Glu Val Ile Asp Ser Asp Phe Glu Thr Arg Asn ttc tgg atc ggg ctc acc tac aag acc gcc aag gac tcc ttc cgc tgg Phe Trp Ile Gly Leu Thr Tyr Lys Thr Ala Lys Asp Ser Phe Arg Trp gcc aca ggg gag cac cag gcc ttc acc agt ttt gcc ttt ggg cag cct Ala Thr Gly Glu His Gln Ala Phe Thr Ser Phe Ala Phe Gly Gln Pro gac aac cac ggg ttt ggc aac tgc gtg gag ctg cag gct tca gct gcc Asp Asn His Gly Phe Gly Asn Cys Val Glu Leu Gln Ala Ser Ala Ala ttc aac tgg aac aac cag cgc tgc aaa acc cga aac cgt tac atc tgc Phe Asn Trp Asn Asn Gln Arg Cys Lys Thr Arg Asn Arg Tyr Ile Cys cag ttt gcc cag gag cac atc tcc cgg tgg ggc cca ggg tcc Gln Phe Ala Gln Glu His Ile Ser Arg Trp Gly Pro Gly Ser

58 /346

tgaggcetga ccacatgget ecetegeetg ecetgggage aceggetetg ettacetgte 1459
egeceacetg tetggaacaa gggecaggtt aagaccacat geeteatgte caaagaggte 1519
teagacettg cacaatgeea gaagttggge agaggagge agggaggeea gtgagggeea 1579
gggagtgagt gttagaagaa getggggeee ttegeetget tttgattggg aagatggget 1639
teaattagat ggeaaaggag aggacacege eagtggteea aaaaggetge teetetteeae 1699
etggeecaga eeetgtgggg eageggaget teeetgtgge atgaaceeea eagggtatta 1759
aattatgaat eagetg

<210> 26

10 <211> 1372

5

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

15 <222> (327)..(920)

<400> 26

20

25

aactgcccgc agtgcccatg gtggctcgga tgggaggaac caccgcggag ccggggacag 60 ggggagcagg gcagtgctct gctgggtgag gggcacccag ctccagaggc taggtgggcg 120 tcgctggtgg gtggactcct gggcgctgcg cggagccgcg ccggctgggt tagcgcgggc 180 ggggcgctta gtcccacccc cagaggaggc ggaagaggag cccgagcctg gccgcgggct 240 gggccccgcc gcagctccag ctggccggct tggtcctgcg gtcccttctc tgggaggccc 300 gaccccggcc gcgcccagcc cccacc atg cca ccc gcg ggg ctc cgc cgg gcc 353

Met Pro Pro Ala Gly Leu Arg Arg Ala

1 5

gcg ccg ctc acc gca atc gct ctg ttg gtg ctg ggg gct ccc ctg gtg 401